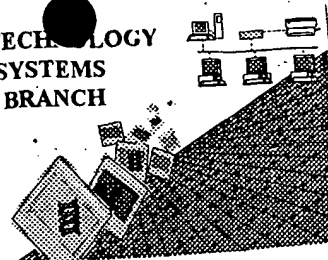


PUT 1/0

BIOTECHNOLOGY
SYSTEMS
BRANCH



#15

CRF Problem Report

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):

Application Serial Number: 10/049,967B
Filing Date: 2/18/02
Date Processed by STIC: 9/10/04

STIC Contact: Mark Spencer: Telephone: 571-272-2510; Fax: 571-273-0221

Nature of Problem:

The CRF (was):

- ☐ (circle one) Damaged or Unreadable (for Unreadable, see attached)
☐ Blank (no files on CRF) (see attached)
☐ Empty file (filename present, but no bytes in file) (see attached)
☐ Virus-infected. Virus name: _____ The STIC will not process the CRF.
☒ Not saved in ASCII text (see attached)
☐ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.
☐ Did not contain a Sequence Listing. (see attached sample)
☐ Other: _____

**PLEASE USE THE CHECKER VERSION 4.2 PROGRAM TO REDUCE ERRORS.
SEE BELOW FOR ADDRESS:**

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/19/04

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The submitted file was saved
in "pdf" (Acrobat) format, which is
invalid per Sequence Rules. 1.824 of Sequence
states that sequence listing
file be saved in ASCII
text format.

10/049,967 B 2

Do NOT show line numbers. This is a scanned
version of most
of previous
submissions.
delete alphabetical headings

delete

3 <110> ~~APPLICANT: Imperial College Innovations Limited~~
5 <120> ~~TITLE OF INVENTION: Methods of treatment~~ This title
7 <130> ~~FILE REFERENCE: ICOY/P23294PC~~ does not
2-> 9 <140> ~~CURRENT APPLICATION NUMBER: US/10/049,967~~ match
2-> 10 <141> ~~CURRENT FILING DATE: 2002-02-18~~ invention title
12 <160> ~~NUMBER OF SEQ ID NOS: 32~~ 28 shown in submitted in application
14 <170> ~~SOFTWARE: PatentIn Ver. 2.0~~ file (p.6) (see bottom of
16 <210> ~~SEQ ID NO: 1~~
17 <211> ~~LENGTH: 25~~
18 <212> ~~TYPE: DNA~~
19 <213> ~~ORGANISM: Artificial Sequence~~
21 <220>
22 <223> ~~OTHER INFORMATION: Description of Artificial Sequence: pcr primer~~
24 <400> ~~SEQUENCE: 1~~
25 agacggatag catggccgag gacgc 25
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33 <223> ~~OTHER INFORMATION: Description of Artificial Sequence: pcr primer~~
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60 <210> ~~SEQ ID NO: 5~~
61 <211> ~~LENGTH: 32~~
62 <212> ~~TYPE: DNA~~
63 <213> ~~ORGANISM: Artificial Sequence~~

Please
consult
Sequence
Rules
for valid
format.

Another invention title given in application: "Isoforms of Snare
Molecules and the
Uses thereof in Modulation
of Cellular Exocytosis"

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3

63 <220>
66 <223> OTHER INFORMATION: Description of Artificial Sequence: pcr primer
68 <400> SEQUENCE: 5
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94 <211> LENGTH: 27
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127 <211> LENGTH: 35
128 <212> TYPE: DNA
129 <213> ORGANISM: Artificial Sequence
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Delete

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~~no~~ no page has allowed

primer?

132 <223> ~~OTHER INFORMATION:~~ Description of Artificial Sequence: pcr ~~primer~~
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line
nos.

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239 <211> LENGTH: 10 codes between
240 <212> TYPE: PRT amino acid
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247 Gln Thr Ala Thr Lys Met Leu Gly Ser Gly
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264 <210> SEQ ID NO: 23 the amino acids under
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266 <212> TYPE: PRT acids
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337 <400> SEQUENCE: 28
338 Gln Thr Gln Thr

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line nos. →

last sequence in submitted file